

Tuesday, March 02, 2004 9:47 AM

Wilbur-Lipman DNA Alignment

Ktuple: 3; Gap Penalty: 3; Window: 20

Seq1(1>379) 491c7 seq 52 (379<-1)	Seq2(1>1518) 491c7 seq 305 (619>997)	Similarity Index 100.0	Gap Number 0	Gap Length 0	Consensus Length 379
v370	v360	v350	v340	v330	v320
ACCCCTGGACTCATACCTGAAAGCAGTGTTCACACCTTAGCAAAATCTCCAACCAGCGCATGAACAATTTC					v310
ACCCCTGGACTCATACCTGAAAGCAGTGTTCACACCTTAGCAAAATCTCCAACCAGCGCATGAACAATTTC					
ACCCCTGGACTCATACCTGAAAGCAGTGTTCACACCTTAGCAAAATCTCCAACCAGCGCATGAACAATTTC					
^620	^630	^640	^650	^660	^670
v300	v290	v280	v270	v260	v250
TACATCACAAACGACCTGGTTTCAAATTCAAGCTCTCAAGGCCAAATCTTTCTAAATTAAACCAAGAACT					v240
TACATCACAAACGACCTGGTTTCAAATTCAAGCTCTCAAGGCCAAATCTTTCTAAATTAAACCAAGAACT					
TACATCACAAACGACCTGGTTTCAAATTCAAGCTCTCAAGGCCAAATCTTTCTAAATTAAACCAAGAACT					
^690	^700	^710	^720	^730	^740
v230	v220	v210	v200	v190	v180
TCATCAGTTCACAGAGAAAGTAATCCAGGACCGGAAGGAGTCTCTTAAGGATAAGCTAAAACAAGATACT					v170
TCATCAGTTCACAGAGAAAGTAATCCAGGACCGGAAGGAGTCTCTTAAGGATAAGCTAAAACAAGATACT					
TCATCAGTTCACAGAGAAAGTAATCCAGGACCGGAAGGAGTCTCTTAAGGATAAGCTAAAACAAGATACT					
^760	^770	^780	^790	^800	^810
v160	v150	v140	v130	v120	v110
ACTCAGAAAAGGCCTGGGATTTCTGGACATACTTTGAGTGCCAAAGCGAAAACACCAAAGATTCT					v100
ACTCAGAAAAGGCCTGGGATTTCTGGACATACTTTGAGTGCCAAAGCGAAAACACCAAAGATTCT					
ACTCAGAAAAGGCCTGGGATTTCTGGACATACTTTGAGTGCCAAAGCGAAAACACCAAAGATTCT					
^830	^840	^850	^860	^870	^880
v90	v80	v70	v60	v50	v40
CTGAAGCAGATCTCAGGCTGAAGTAAAAACGTTCATGTTGCAGGACATGACACCACATCCAGTGCTAT					v30
CTGAAGCAGATCTCAGGCTGAAGTAAAAACGTTCATGTTGCAGGACATGACACCACATCCAGTGCTAT					
CTGAAGCAGATCTCAGGCTGAAGTAAAAACGTTCATGTTGCAGGACATGACACCACATCCAGTGCTAT					
^900	^910	^920	^930	^940	^950
v20	v10				
CTCCTGGATCCTTACTGCTTGGCAAAGT					
CTCCTGGATCCTTACTGCTTGGCAAAGT					
CTCCTGGATCCTTACTGCTTGGCAAAGT					
^970	^980	^990			

Figure 1

Page 1



Figure 2

B854P Extended Breast Panel

